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SEQUENCE LISTING

(1)	GENERAL	INFORMATION:

- (I) APPLICANT: The Scripps Research Institute
- (ii) TITLE OF INVENTION: ENZYMATIC DNA MOLECULES
 - (iii) NUMBER OF SEQUENCES: 101
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: The Scripps Research Institute
 - (B) STREET: 10666 North Torrey Pines Road, TPC-8
- 10 (C) CITY: La Jolla
 - (D) STATE: California
 - (E) COUNTRY: United States
 - (F) ZIP: 92037
- 15 (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/US95/
 - (B) FILING DATE: 01-DEC-1995
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/472,194
 - (B) FILING DATE: 07-JUN-1995
- 30 (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/349,023
 - (B) FILING DATE: 02-DEC-1994
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Logan, April C.
 - (B) REGISTRATION NUMBER: 33,950
 - (C) REFERENCE/DOCKET NUMBER: 463.2 PC
 - (ix) TELECOMMUNICATION INFORMATION:
- **40** (A) TELEPHONE: (619) 554-2937

(B) TELEFAX: (619) 554-6312

(2) INFORMATION FOR SEQ ID NO:1:

5

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

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- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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CGGTAAGCTT GGCAC

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- (2) INFORMATION FOR SEQ ID NO:2:
- 20 (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

25

- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: misc_difference
 - (B) LOCATION: replace(8, "")
 - (D) OTHER INFORMATION: /standard_name= "ADENOSINE RIBONUCLEOTIDE"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

	TCACTATNAG GAAGAGATGG	20
5	(2) INFORMATION FOR SEQ ID NO:3:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 38 base pairs	
10	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
	ACACATCTCT GAAGTAGCGC CGCCGTATAG TGACGCTA	38
20	(2) INFORMATION FOR SEQ ID NO:4:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 80 base pairs(B) TYPE: nucleic acid	
25	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	·

NNNNGTCGC CATCTCTTCC

1	2	INFORMATION	FOR	SEO	TD	NO.5
١	٠ ،	THEOMETICAL	LON	עייי	10	140.0

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 28 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

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(ix) FEATURE:

- (A) NAME/KEY: misc feature
- (B) LOCATION: 28

15 (D) OTHER INFORMATION: /standard_name= "2'3' CYCLIC PHOSPHATE"

(ix) FEATURE:

- (A) NAME/KEY: misc_difference
- (B) LOCATION: replace(28, "")
- (D) OTHER INFORMATION: /standard_name= "ADENOSINE RIBONUCLEOTIDE"

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GGGACGAATT CTAATACGAC TCACTATN

(2) INFORMATION FOR SEQ ID NO:6:

30

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
- 35 (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
- 40 (ix) FEATURE:

(ix) FEATURE:

		(A) NAME/KEY: misc_difference(B) LOCATION: replace(28, "")(D) OTHER INFORMATION: /standard_name= "ADENOSINE RIBONUCLEOTIDE"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
GGGA	CGAA!	TT CTAATACGAC TCACTATN	28
(2)	INFO	RMATION FOR SEQ ID NO:7:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 19 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(ix)	FEATURE:	
		(A) NAME/KEY: misc_difference	
		(B) LOCATION: replace(8, "")	
		(D) OTHER INFORMATION: /standard_name= "ADENOSINE	
		RIBONUCLEOTIDE"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
TCAC	CTATN	GG AAGAGATGG	19
(2)	INFO	RMATION FOR SEQ ID NO:8:	
	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 8 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	

	(A) NAME/KEY: misc_difference	
	(B) LOCATION: replace(8, "")	
	(D) OTHER INFORMATION: /standard_name= "ADENOSINE	•
	NUCLEOTIDE"	
5		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
	TCACTATN	8
10	(2) INFORMATION FOR SEQ ID NO:9:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 30 base pairs	
	(B) TYPE: nucleic acid	
15	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
20	(XI) SECOUNCE DESCRIPTION. SEC ID NO. 3.	
	CCATCTCTTC CTATAGTGAG TCCGGCTGCA	30
	(2) INFORMATION FOR SEQ ID NO:10:	
25		
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 15 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
30	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
35		
	GTGCCAAGCT TACCG	15
	(a) Turnery Ten one of the Valla	
	(2) INFORMATION FOR SEQ ID NO:11:	
4.0	/÷ \ GEOMENOS GUADAGERATOS	
40	(i) SEQUENCE CHARACTERISTICS:	

		(A)	LENGTH: 43 base pairs	
		(B)	TYPE: nucleic acid	
		(C)	STRANDEDNESS: single	
		(D)	TOPOLOGY: linear	
(ii)	MOLE	CULE TYPE: DNA (genomic)	
(xi)	SEQUE	ENCE DESCRIPTION: SEQ ID NO:11:	
CTGCA	GAA?	гт стл	AATACGAC TCACTATAGG AAGAGATGGC GAC	43
(2) I	NFOF	RMATIO	ON FOR SEQ ID NO:12:	
	(i)	SEQUI	ENCE CHARACTERISTICS:	
		(A)	LENGTH: 19 base pairs	
		(B)	TYPE: nucleic acid	
		(C)	STRANDEDNESS: single	
		(D)	TOPOLOGY: linear	
(ii)	MOLE	CULE TYPE: DNA (genomic)	
(ix)	FEAT	JRE:	
		(A)	NAME/KEY: misc_difference	
		(B)	LOCATION: replace(8, "")	
		(D)	OTHER INFORMATION: /standard_name= "ADENOSINE	
			RIBONUCLEOTIDE"	
((xi)	SEQU	ENCE DESCRIPTION: SEQ ID NO:12:	
TCACI	CATN	gg aa	GAGATGG	19
(2) I	NFO:	RMATI	ON FOR SEQ ID NO:13:	
	(i)	SEQU	ENCE CHARACTERISTICS:	
		(A)	LENGTH: 43 base pairs	
		(B)	TYPE: nucleic acid	
		(C)	STRANDEDNESS: single	
		(D)	TOPOLOGY: linear	

(ii) MOLECULE TYPE: DNA (genomic)

	(ix) FEATURE:	
	(A) NAME/KEY: misc_difference	
	(B) LOCATION: replace(28, "")	
	(D) OTHER INFORMATION: /standard_name= "ADENOSINE	
5	RIBONUCLEOTIDE"	
	·	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
	GGGACGAATT CTAATACGAC TCACTATNGG AAGAGATGGC GAC	43
10	(2) INFORMATION FOR SEQ ID NO:14:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
15	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
	TCACACATCT CTGAAGTAGC GCCGCCGTAT GTGACGCTAG GGGTTCGCCT	50
	TOACACATOT CIGAAGIAGE GCCCCCGIAT GTGACGCIAG GCCTTCCCCT	00
25	(2) INFORMATION FOR SEQ ID NO:15:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
30	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
	GGGGGAACG CCGTAACAAG CTCTGAACTA GCGGTTGCGA TATAGTCGTA	50
	(2) INFORMATION FOR SEQ ID NO:16:	
40		

(A) LENGTH: 50 base pairs	
(,	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
CGGGACTCCG TAGCCCATTG CTTTTTGCAG CGTCAACGAA TAGCGTATTA	50
(2) INFORMATION FOR SEQ ID NO:17:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 50 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(x1) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
CCACCATGTC TTCTCGAGCC GAACCGATAG TTACGTCATA CCTCCCGTAT	50
(2) INFORMATION FOR SEQ ID NO:18:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 50 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
GCCAGATTGC TGCTACCAGC GGTACGAAAT AGTGAAGTGT TCGTGACTAT	50
(2) INFORMATION FOR SEQ ID NO:19:	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16: CGGGACTCCG TAGCCCATTG CTTTTTGCAG CGTCAACGAA TAGCGTATTA (2) INFORMATION FOR SEQ ID NO:17: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17: CCACCATGTC TTCTCGAGCC GAACCGATAG TTACGTCATA CCTCCCGTAT (2) INFORMATION FOR SEQ ID NO:18: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18: GCCAGATTGC TGCTACCAGC GGTACGAAAT AGTGAAGTGT TCGTGACTAT

	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
5	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
10	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
10	ATAGGCCATG CTTTGGCTAG CGGCACCGTA TAGTGTACCT GCCCTTATCG	50
	(2) INFORMATION FOR SEQ ID NO:20:	
15	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
20	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
25	TCTGCTCTCC TCTATTCTAG CAGTGCAGCG AAATATGTCG AATAGTCGGT	50
	(2) INFORMATION FOR SEQ ID NO:21:	
	(i) SEQUENCE CHARACTERISTICS:	
30	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
35	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
40	TTGCCCAGCA TAGTCGGCAG ACGTGGTGTT AGCGACACGA TAGGCCCGGT	50

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	(2) INFORMATION FOR SEQ ID NO:22:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
5	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
10	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
	TTGCTAGCTC GGCTGAACTT CTGTAGCGCA ACCGAAATAG TGAGGCTTGA	50
15	(2) INFORMATION FOR SEQ ID NO:23:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 107 base pairs	
	(B) TYPE: nucleic acid	
20	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
25	(ix) FEATURE:	
	(A) NAME/KEY: misc_difference	
	(B) LOCATION: replace(28, "")	
	(D) OTHER INFORMATION: /standard_name= "ADENOSINE	
	RIBONUCLEOTIDE"	
30	/label= rA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
	GGGACGAATT CTAATACGAC TCACTATNGG AAGAGATGGC GACATCTCNN NNNNNNNNN	60
35	NNNNNNNNN NNNNNNNN NNNNNNNNGT GACGGTAAGC TTGGCAC	107
	/O/ TNEODWRETON FOR CEO TO NO.O.	

(2) INFORMATION FOR SEQ ID NO:24:

(i) SEQUENCE CHARACTERISTICS:

40 (A) LENGTH: 49 base pairs

	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
5	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
10	CCGCCCACCT CTTTTACGAG CCTGTACGAA ATAGTGCTCT TGTTAGTAT	49
10	(2) INFORMATION FOR SEQ ID NO:25:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 48 base pairs	
15	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
•	(ii) MOLECULE TYPE: DNA (genomic)	
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
	TCTCTTCAGC GATGCACGCT TGTTTTAATG TTGCACCCAT GTTAGTGA	48
25	(2) INFORMATION FOR SEQ ID NO:26:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 46 base pairs	
	(B) TYPE: nucleic acid	
30	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
	TCTCATCAGC GATTGAACCA CTTGGTGGAC AGACCCATGT TAGTGA	46
	(2) INFORMATION FOR SEQ ID NO:27:	
40		

5	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 49 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
10	CCGCCCACCT CTTTTACGAG CCTGTACGAA ATAGTGTTCT TGTTAGTAT	49
	(2) INFORMATION FOR SEQ ID NO:28:	
15	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 49 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
20		
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
25	CCGCCCACCT CTTTTACGAG CCTGTACGAA ATAGTGCTCT CGTTAGTAT	49
	(2) INFORMATION FOR SEQ ID NO:29:	
	(i) SEQUENCE CHARACTERISTICS:	
30	(A) LENGTH: 48 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
35	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
	TCTCAGACTT AGTCCATCAC ACTCTGTGCA TATGCCTGCT TGATGTGA	48
40		

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	(2) INFORMATION FOR SEQ ID NO:30:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 42 base pairs	
5	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
10	(ii) MOLECULE TYPE: DNA (genomic)	
10	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
	CTCTCATCTG CTAGCACGCT CGAATAGTGT CAGTCGATGT GA	42
15	(2) INFORMATION FOR SEQ ID NO:31:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 40 base pairs	
	(B) TYPE: nucleic acid	
20	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
25	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
	TACAGCGATT CACCCTTGTT TAAGGGTTAC ACCCATGTTA	40
30	(2) INFORMATION FOR SEQ ID NO:32:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 40 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
35	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

ATCAGCGATT AACGCTTGTT TCAATGTTAC ACCCATGTTA 40 (2) INFORMATION FOR SEQ ID NO:33: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: 40 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33: 15 TTCAGCGATT AACGCTTATT TTAGCGTTAC ACCCATGTTA 40 (2) INFORMATION FOR SEQ ID NO:34: (i) SEQUENCE CHARACTERISTICS: 20 (A) LENGTH: 40 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 25 (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34: ATCAGCGATT CACCCTTGTT TTAAGGTTGC ACCCATGTTA 40 30 (2) INFORMATION FOR SEQ ID NO:35: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 40 base pairs 35 (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) 40

	-114-	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:	
	ATCAGCGATT CACCCTTGTT TAAGCGTTAC ACCCATGTTG	40
5	(2) INFORMATION FOR SEQ ID NO:36:	
•	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 40 base pairs	
	(B) TYPE: nucleic acid	
. 10	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:	
	ATCAGCGATT CACCCTTGTT TTAAGGTTAC ACCCATGTTA	40
	(2) INFORMATION FOR SEQ ID NO:37:	
20		
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 40 base pairs	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
25		
23	(b) Toroboot. Iffical	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:	
30	ATCAGCGATT AACGCTTATT TTAGCGTTAC ACCCATGTTA	40
	(2) INFORMATION FOR SEQ ID NO:38:	
35	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 40 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	

(D) TOPOLOGY: linear

	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:	
5	ATCAGCGATT AACGCTTGTT TTAGTGTTGC ACCCATGTTA	40
	(2) INFORMATION FOR SEQ ID NO:39:	
	(i) SEQUENCE CHARACTERISTICS:	
10	(A) LENGTH: 40 base pairs	
10	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
15	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
20	ATCAGCGATT AACGCTTATT TTAGCATTAC ACCCATGTTA	40
20	(2) INFORMATION FOR SEQ ID NO:40:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 10 base pairs	
25	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
30	(ii) MOLECULE TYPE: DNA (genomic)	
30	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
	GCCATGCTTT	10
35	(2) INFORMATION FOR SEQ ID NO:41:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 10 base pairs	
	(B) TYPE: nucleic acid	
40	(C) STRANDEDNESS: single	

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		-116-	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
5	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:	
	CTCTATTT	CT	10
	(2) INFO	RMATION FOR SEQ ID NO:42:	
10	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 12 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
15		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
20	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:	
20	TATGTGAC	GC TA	12
	(2) INFO	RMATION FOR SEQ ID NO:43:	
25	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 10 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
30			
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:	
35	TATAGTCG	TA	10

(2) INFORMATION FOR SEQ ID NO:44:

(i) SEQUENCE CHARACTERISTICS:

40 (A) LENGTH: 11 base pairs

	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
5	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:	
	ATAGCGTATT A	11
10	(2) INFORMATION FOR SEQ ID NO:45:	
15	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 13 base pairs(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
20	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:	
	ATAGTTACGT CAT	13
25	(2) INFORMATION FOR SEQ ID NO:46:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs	
30	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
35	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:	
	AATAGTGAAG TGTT	14
	(2) INFORMATION FOR SEQ ID NO:47:	

(i) SEQUENCE CHARACTERISTICS:

WO 98/49346

	(A) LENGTH: 11 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
5	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
10	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:	
10	ATAGGCCCGG T	11
	(2) INFORMATION FOR SEQ ID NO:48:	
15	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 14 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
20		
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:	
25	AATAGTGAGG CTTG	14
	(2) INFORMATION FOR SEQ ID NO:49:	
	(i) SEQUENCE CHARACTERISTICS:	
30	(A) LENGTH: 12 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
35	(ii) MOLECULE TYPE: RNA (genomic)	
	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:	
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GUAACUAGAG AU	12
(2) INFORMATION FOR SEQ ID NO:50:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 98 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: NO	
(ix) FEATURE:	
(A) NAME/KEY: misc_feature	
(B) LOCATION: 718	
(D) OTHER INFORMATION: /note= "Position 7-18 is RNA; the	

20 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

GGAAAAGUAA CUAGAGAUGG AAGAGATGGC GACNNNNNN NNNNNNNNN NNNNNNNNNN 60 NNNNNNNNN NNNNNNNNN NNNCGGTAAG CTTGGCAC 98

remainder of the sequence is DNA."

25 (2) INFORMATION FOR SEQ ID NO:51:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 99 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- 35 (iv) ANTI-SENSE: NO
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..24
 - (D) OTHER INFORMATION: /note= "Positions 1-24 is RNA; the remainder of the sequence is DNA."

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:

	GGAAAAGUA ACUAGAGAUG GAAGAGATGG CGACNNNNNN NNNNNNNNNN NNNNNNNNNNNNNNN	60 99
5	(2) INFORMATION FOR SEQ ID NO:52:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
10	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
15	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:	
20	CCAATAGTGC TACTGTGTAT CTCAATGCTG GAAACACGGG TTATCTCCCG	50
	(2) INFORMATION FOR SEQ ID NO:53:	
	(i) SEQUENCE CHARACTERISTICS:	
25	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
30	(ii) MOLECULE TYPE: DNA (genomic)	
	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:	•
35	CCAAAACAGT GGAGCATTAT ATCTACTCCA CAAAGACCAC TTTTCTCCCG	50
	(2) INFORMATION FOR SEQ ID NO:54:	
		-
40	(i) SEQUENCE CHARACTERISTICS:	

	(A) LENGTH: 50 base pairs	•
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:54:	
ATCCGTACT	TA GCATGCAGAC AGTCTGTCTG CTTTTTCATT ACTCACTCCC	50
(2) INFO	RMATION FOR SEQ ID NO:55:	
/÷ \	CECUENCE CURRECTED CONT.CC.	
(1)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 49 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
	HYPOTHETICAL: NO	
•	ANTI-SENSE: NO	
	SEQUENCE DESCRIPTION: SEQ ID NO:55:	
(7		
CAATTCAT	GA TGACCAACTC TGTCAACACG CGAACTTTTA ACACTGGCA	49
(2) INFO	RMATION FOR SEQ ID NO:56:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:

CTTCCACCTT CCGAGCCGGA CGAAGTTACT TTTTATCACA CTACGTATTG 50 (2) INFORMATION FOR SEQ ID NO:57: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:57: GGCAAGAGAT GGCATATATT CAGGTAACTG TGGAGATACC CTGTCTGCCA 50 (2) INFORMATION FOR SEQ ID NO:58: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:58: CTAGACCATT CACGTTTACC AAGCTATGGT AAGAACTAGA ATCACGCGTA 50 35 (2) INFORMATION FOR SEQ ID NO:59: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid 40 (C) STRANDEDNESS: single

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- 5 (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:

CGTACACGTG GAAAAGCTAT AAGTCAAGTT CTCATCATGT ACCTGACCGC 50

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- (2) INFORMATION FOR SEQ ID NO:60:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 50 base pairs
- 15 (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
- 20 (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:

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CAGTGATACA TGAGTGCACC GCTACGACTA AGTCTGTAAC TTATTCTACC 50

- (2) INFORMATION FOR SEQ ID NO:61:
- 30 (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 50 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- 40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:

	ACCGAATTAA ACTACCGAAT AGTGTGGTTT CTATGCTTCT TCTTCCCTGA	50
	(2) INFORMATION FOR SEQ ID NO:62:	
5	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 50 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
10		
	(ii) MOLECULE TYPE: DNA (genomic)	
	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
15	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:	
	CAGGTAGATA TAATGCGTCA CCGTGCTTAC ACTCGTTTTA TTAGTATGTC	50
	(2) INFORMATION FOR SEQ ID NO:63:	
20	(i) CHOURNER CHARACTERICA.	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 49 base pairs (B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
25	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
30		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:	
	CCCTACAACA CCACTGGGCC CAATTAGATT AACGCTATTT TATAACTCG	49
35	(2) INFORMATION FOR SEQ ID NO:64:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 49 base pairs	
	(B) TYPE: nucleic acid	
40	(C) STRANDEDNESS: single	

(D)	TOPOLOGY:	linear
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1441	MOTECUTE	TYPE: DNA	/ = = = = = = = = = = = = = = = = = = =
(111)	MODERCULE	TOPE: DNA	(denomic)

- (iii) HYPOTHETICAL: NO
- 5 (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:

CCAAACGGTT ATAAGACTGA AAACTCAATC AATAGCCCAA TCCTCGCCC 49

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(2) INFORMATION FOR SEQ ID NO:65:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 50 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:

25 CACATGTATA CCTAAGAAAT TGGTCCCGTA GACGTCACAG ACTTACGCCA 50

(2) INFORMATION FOR SEQ ID NO:66:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 50 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

35 (ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:

CACAACGAAA ACAATCTTCC TTGGCATACT GGGGAGAAAG TCTGTTGTCC 50 (2) INFORMATION FOR SEQ ID NO:67: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:67: CACACGAACA TGTCCATTAA ATGGCATTCC GTTTTTCGTT CTACATATGC 50 20 (2) INFORMATION FOR SEQ ID NO:68: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 49 base pairs (B) TYPE: nucleic acid 25 (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO 30 (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:68: CAGAACGAGG GTCTTGTAAG ACTACACCTC CTCAGTGACA ATAATCCTG 49 35 (2) INFORMATION FOR SEQ ID NO:69: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 49 base pairs 40 (B) TYPE: nucleic acid

		<pre>(C) STRANDEDNESS: single (D) TOPOLOGY: linear</pre>	
5	(iii)	MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO ANTI-SENSE: NO	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:69:	
10	CACTACAG	CC TGATATATAT GAAGAACAGG CAACAAGCTT ATGCACTGG	49
	(2) INFO	RMATION FOR SEQ ID NO:70:	
	(i)	SEQUENCE CHARACTERISTICS:	
15		(A) LENGTH: 50 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
20	(ii)	MOLECULE TYPE: DNA (genomic)	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
25	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:70:	
	GGGTACAT	TT ATGATTCTCT TATAAAGAGA ATATCGTACT CTTTTCCCCA	50
	(2) INFO	RMATION FOR SEQ ID NO:71:	
30	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 49 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
35			
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:

CCAAAGTACA TTCCAACCCC TTATACGTGA AACTTCCAGT AGTTTCCTA 49 (2) INFORMATION FOR SEQ ID NO:72: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:72: 50 CTTGAAGATC CTCATAAGAC GATTAAACAA TCCACTGGAT ATAATCCGGA (2) INFORMATION FOR SEQ ID NO:73: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:73: CGAATAGTGT CCATGATTAC ACCAATAACT GCCTGCCTAT CATGTTTATG 50 35 (2) INFORMATION FOR SEQ ID NO:74: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid 40 (C) STRANDEDNESS: single

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-129-(D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:74: CCAAGAGAGT ATCGGATACA CTTGGAACAT AGCTAACTCG AACTGTACCA 50 (2) INFORMATION FOR SEQ ID NO:75: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:75: CCACTGATAA ATAGGTAACT GTCTCATATC TGCCAATCAT ATGCCGTA 48

(2) INFORMATION FOR SEQ ID NO:76:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 50 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:

CCCAAATTAT AAACAATTTA ACACAAGCAA AAGGAGGTTC ATTGCTCCGC 50 (2) INFORMATION FOR SEQ ID NO:77: 5 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:77: CAATAAACTG GTGCTAAACC TAATACCTTG TATCCAAGTT ATCCTCCCCC 50 (2) INFORMATION FOR SEQ ID NO:78: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:78: CCGAATGACA TCCGTAGTGG AACCTTGCTT TTGACACTAA GAAGCTACAC 50 35 (2) INFORMATION FOR SEQ ID NO:79: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) T	OPOLOGY:	linear
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(ii)	MOLECULE	TYPE:	DNA	(genomic)
· — — /				, 9 0 0 2 . ,

- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:

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(2) INFORMATION FOR SEQ ID NO:80:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 50 base pairs
- 15 (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
- 20 (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:80:

25 CAGAACAAG ATCAGTAGCT AAACATATGG TACAAACATA CCATCTCGCA 50

- (2) INFORMATION FOR SEQ ID NO:81:
 - (i) SEQUENCE CHARACTERISTICS:
- 30 (A) LENGTH: 49 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- 35 (ii) MOLECULE TYPE: DNA (genomic)
 - (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
- 40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:

CCTTTAGTTA GGCTAGCTAC AACGATTTTT CCCTGCTTGG CAACGACAC 49 (2) INFORMATION FOR SEQ ID NO:82: 5 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:82: CTCCCTACGT TACACCAGCG GTACGAATTT TCCACGAGAG GTAATCCGCA 50 (2) INFORMATION FOR SEQ ID NO:83: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:83: CGGCACCTCT AGTTAGACAC TCCGGAATTT TTCCCC 36 35 (2) INFORMATION FOR SEQ ID NO:84: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 49 base pairs (B) TYPE: nucleic acid 40 (C) STRANDEDNESS: single

(D)	TOPO	LOGY:	linear
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- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- 5 (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:

CGGCACCTCT AGTTAGACAC TCCGGAATTT TAGCCTACCA TAGTCCGGT 49

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- (2) INFORMATION FOR SEQ ID NO:85:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 47 base pairs
- 15 (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
- 20 (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:85:
- 25 CCCTTTGGTT AGGCTAGCTA CAACGATTTT TCCCTGCTTG AATTGTA 47
 - (2) INFORMATION FOR SEQ ID NO:86:
 - (i) SEQUENCE CHARACTERISTICS:
- 30 (A) LENGTH: 51 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- 35 (ii) MOLECULE TYPE: DNA (genomic)
 - (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:

CCCTTTGGTT AGGCTAGCTA CAACGATTTT TCCCTGCTTG ACCTGTTACG A 51 (2) INFORMATION FOR SEQ ID NO:87: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:87: CCTTTAGTTA GGCTAGCTAC AACGATTTTT CCCTGCTTGG AACGACAC 48 (2) INFORMATION FOR SEQ ID NO:88: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO 30 (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:88: CATGGCTTAA TCATCCTCAA TAGAAGACTA CAAGTCGAAT ATGTCCCCCC 50 35 (2) INFORMATION FOR SEQ ID NO:89: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs 40 (B) TYPE: nucleic acid

		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
5	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:89:	
10	CAACAGAG	CG AGTATCACCC CCTGTCAATA GTCGTATGAA ACATTGGGCC	50
	(2) INFO	RMATION FOR SEQ ID NO:90:	
	(i)	SEQUENCE CHARACTERISTICS:	
15		(A) LENGTH: 49 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
20	(ii)	MOLECULE TYPE: DNA (genomic)	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
25	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:90:	
23	TACCGACA	AG GGGAATTAAA AGCTAGCTGG TTATGCAACC CTTTTCGCA	49
	(2) INFO	RMATION FOR SEQ ID NO:91:	
30	(i)	SEQUENCE CHARACTERISTICS:	
		(A) LENGTH: 49 base pairs	
		(B) TYPE: nucleic acid	
		(C) STRANDEDNESS: single	
		(D) TOPOLOGY: linear	
35			
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
40	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:91:	

CTCGAAACAG TGATATTCTG AACAAACGGG TACTACGTGT TCAGCCCCC 49 (2) INFORMATION FOR SEQ ID NO:92: 5 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:92: CCAATAACGT AACCCGGTTA GATAAGCACT TAGCTAAGAT GTTTATCCTG 50 (2) INFORMATION FOR SEQ ID NO:93: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:93: CAATACAATC GGTACGAATC CAGAAACATA ACGTTGTTTC AGAATGGTCC 50 35 (2) INFORMATION FOR SEQ ID NO:94: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid 40

(C) STRANDEDNESS: single

(D)	TOPOLOGY:	linear
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11	. 1. 1	MOTECOTE	TYPEI	UNA	(denomic)

- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:94:

GCAACAACAA GAACCAAGTT ACATACACGT TCATCTATAC TGAACCCCCA 50

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(2) INFORMATION FOR SEQ ID NO:95:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 50 base pairs
- 15 (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
- 20 (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:95:

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CCTTTGAGTT CCTAAATGCC GCACGGTAAG CTTGGCACAC TTTGACTGTA 50

- (2) INFORMATION FOR SEQ ID NO:96:
- 30 (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 49 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- 40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:96:

CAAAGATCTC ACTTTGGAAA TGCGAAATAT GTATATTCGC CCTGTCTGC 49 (2) INFORMATION FOR SEQ ID NO:97: 5 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 10 (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:97: CCACGTAGAA TTATCTGATT TATAACATAA CGCAGGATAA CTCTCGCCCA 50 (2) INFORMATION FOR SEQ ID NO:98: 20 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single 25 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:98: CACAAGAAG TGTCGTCTCC AGATATTTGA GTACAAGGAA CTACGCCC 48 35 (2) INFORMATION FOR SEQ ID NO:99: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid 40 (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:99:

CATGAAGAAA TAGGACATTC TACAGGCTGG ACCGTTACTA TGCCTGTAGG 50

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(2) INFORMATION FOR SEQ ID NO:100:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 46 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:100:

25 CATAGGATAA TCATGGCGAT GCTTATGACG TGTACATCTA TACCTT 46

(2) INFORMATION FOR SEQ ID NO:101:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 50 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

35 (ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:101: